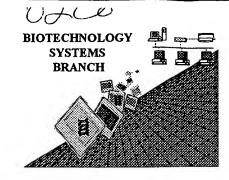
RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following CRF diskette:

Application Serial Number: 09/454,651Art Unit / Team No.: 0/16Date Processed by STIC: 12/20/99

THE ATTACHED PRINTOUT EXPLAINS THE ERRORS DETECTED.

PLEASE BE SURE TO FORWARD THIS INFORMATION TO THE APPLICANTS BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANTS ALONG WITH A NOTICE TO COMPLY or,
- 2) CALLING APPLICANTS AND FAXING THEM A COPY OF THE PRINTOUT WITH A NOTICE TO COMPLY

THIS WILL INSURE THAT THE NEXT SUBMISSION RECEIVED FROM THEM WILL BE ERROR FREE.

IF YOU HAVE ANY FURTHER QUESTIONS, PLEASE CALL:

MARK SPENCER 703-308-4212

RAW SEQUENCE LISTING PATENT APPLICATION US/09/454,651

DATE: 12/20/1999 TIME: 16:47:48

INPUT SET: S34294.raw

This Raw Listing contains the General Information Section and those Sequences containing ERRORS.

	1		SEQUENCE LISTING	Does Not Comply
•	2	(1)	General Information:	Corrected Diskette Needed
	3 4	(1)	General información:	
	5		(i) APPLICANT: Linsley, Peter S.	
	6		Ledbetter, Jeffrey A.	
	7		Damle, Nitin K.	
	8		Brady, William	
	9		Wallace, Philip M.	
	10			
	11		(ii) TITLE OF THE INVENTION: CTLA4/CD28Ig HYBRID F	USION
	12		PROTEINS AND USES THEREOF	
	13		\sim	
>	14		(iii) NUMBER OF SEQUENCES: (22) pur p. 3	•
	15			
	16		(iv) CORRESPONDENCE ADDRESS:	
	17		(A) ADDRESSEE: Merchant & Gould	
	18		(B) STREET: 11150 Santa Monica Boulevard, Suite	400
	19		(C) CITY: Los Angeles	
	20		(D) STATE: CA	
	21		(E) COUNTRY: USA	
	22		(F) ZIP: 90025	
	23			
	24		(v) COMPUTER READABLE FORM:	
	25		(A) MEDIUM TYPE: Diskette	
	26		(B) COMPUTER: IBM Compatible	
	27		(C) OPERATING SYSTEM: DOS	
	28		(D) SOFTWARE: FastSEQ Version 2.0	
	29		(- 1) CURRENT ARRY TOURTON BARRY	o under PRIOR APP. DATA:
	30		(vi) CURRENT APPLICATION DATA.	of under PROTOR 1101.
>	31		(A) APPLICATION NUMBER: 08/228,208 / These (A)	OATA!
	32		(B) FILING DATE: (15-APR-1994 (C) CLASSIFICATION:	
	33 34		(C) CLASSIFICATION:	
	35		(vii) PRIOR APPLICATION DATA:	
	36		(A) APPLICATION NUMBER: 08/008,898	
	37		(B) FILING DATE: 22-JAN-1993	
	38		(D) FIDING DATE: 22 ORN 1993	
	39		(A) APPLICATION NUMBER: 07/723,617	
	40		(B) FILING DATE: 27-JUN-1991	
	41		(b) IIIII biili bi ook 1991	
	42			
	43		(viii) ATTORNEY/AGENT INFORMATION:	
	44		(A) NAME: Adriano, Sarah B	
	45		(B) REGISTRATION NUMBER: 34,470	
			• •	

53

RAW SEQUENCE LISTING PATENT APPLICATION US/09/454,651

DATE: 12/20/1999 TIME: 16:47:48

		INPUT SET: S34294.raw
46	(C) REFERENCE/DOCKET NUMBER: 30436-30US01	
47		
48	(ix) TELECOMMUNICATION INFORMATION:	
49	(A) TELEPHONE: 310 445-1140	
50	(B) TELEFAX: 310 445-9031	
51	(C) TELEX:	
52		

ERRORED SEQUENCES FOLLOW:

	334			(2)) INI	FORM	OITA	V FOI	R SE	Q ID	NO:	17:					
	335														,		_ ^
	336		(:	i) SI	EQUE	NCE (CHAR	ACTE	RIST	ICS:		7	77		/	_ / _	2 C
>	337	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 186 amino acids 223 Mourles 3															
	338	(B) TYPE: amino acid															
	339	(C) STRANDEDNESS: unknown															
	340	(D) TOPOLOGY: linear															
	341																
	342		(:	ii) I	MOLE	CULE	TYPI	E: p	rote:	in							
	343																
	344		(2	ki) S	SEQUI	ENCE	DES	CRIP	rion	: SE	Q ID	NO:	17:				
	345																
	346																
	347	Met	Ala	Cys	Leu	Gly	Phe	Gln	Arg	His	Lys	Ala	Gln	Leu	Asn	Leu	Ala
	348	1				5					10					15	
	349	Ala	Pro	Thr	\mathtt{Trp}	Pro	Cys	Thr	Leu	Leu	Phe	Phe	Leu	Leu	Phe	Ile	Pro
	350				20					25					30		
	351	Val	Phe	Cys	Lys	Ala	Met	His	Val	Ala	Gln	Pro	Ala	Val	Val	Leu	Ala
	352			35					40					45			_
	353	Ser	Ser	Arg	Gly	Ile	Ala	Ser	Phe	Val	Cys	Glu	Tyr	Ala	Ser	Pro	Gly
	354		50					55_	_	_			60	_			
	355	_	Ala	Thr	Glu	Val	_	Val	Thr	Val	Leu	_	Gln	Ala	Asp	Ser	
	356	65	_				70	_	_			75	_		_		80
	357	Val	Thr	Glu	Val	_	Ala	Ala	Thr	Tyr		Met	Gly	Asn	Glu		Thr
	358					85		_			90	_	_		_	95	
	359	Phe	Leu	Asp		Ser	Ile	Cys	Thr		Thr	Ser	ser	GIĀ		GIn	Val
	360	_	_		100			_	_	105		_	_,	~ 7	110	_	
	361	Asn	Leu		Ile	GIn	GIY	Leu	_	Ala	Met	Asp	Thr	_	Leu	Tyr	IIe
	362		_	115		_			120					125	a 1	-1-	a 1
	363	Cys	_	vaı	GIu	Leu	Met	Tyr	Pro	Pro	Pro	Tyr		Leu	GIY	тте	GIA
	364	_	130	_,			_	135	-1-	•		~1	140	a			
	365		GTĀ	Thr	GIn	TTE	_	Val	тте	Asp	Pro		Pro	Cys	Pro	Asp	
	366	145	_,	_ ^	_	_	150	_	- 1 -		7	155	a	~1		5 1	160
	367	Asp	Pne	Leu	ьeu	_	тте	Leu	Ата	Ата		ser	ser	GIY	Leu		Pne
	368	_	_		_	165			7		170				•	175	•
	369	Tyr	ser	Pne		ьeu	Thr	Ala	vaı		Leu	ser	гуѕ	Met		гла	гÀг
	370	3	a	Desc	180	mb	mb	61. -	770 7	185	1707	T	Mot	Dwc	190	mb	~ 1
	371	arg	ser		ьeu	ınr	ınr	Gly		TYT	vai	пĀв	Mec		PIO	inr	GIU
	372			195					200					205			

RAW SEQUENCE LISTING PATENT APPLICATION US/09/454,651

DATE: 12/20/1999 TIME: 16:47:49

INPUT SET: S34294.raw

															INF	PUT S	ET: \$342	29 4.ra w
	373	Pro	Glu	Cys	Glu	Lys	Gln	Phe	Gln	${\tt Pro}$	Tyr	Phe	Ile	Pro	Ile	Asn		
	374		210					215					220					
	375																	
	376																	
				- /-				T F01	- CE	. TD	270 - 1							
	418			(2)) INI	FORM	A.I.TOI	N FOI	K SE(מד ג	NO:	19:						
	419					TOTAL A	7113 D	A COURT	3 T C/M '	rae.								
	420		(:			NCE (GTH :/						11	01	10	11-1			
>	421					E: at				1100		10	V 12	~00				
	422					ANDE				m								
	423					OLOG				VII								
	424 425			(D)	TOP	OLOG.	1. 1.	IIICa.										
	425		1.	; ;) ı	MOT.F	CULE	тур	R ∙ mi	rote:	in								
	427		١.	11, 1	·IOHE,	COLL		о. р.										
	428		(-	vi) :	SEOII	ENCE	DES	CRIP	TION	: SE	Q ID	NO:	19:					
	429		(-	,	JLQU.						~							
	430	Met	Thr	Leu	Ara	Leu	Leu	Phe	Leu	Ala	Leu	Asn	Phe	Phe	Ser	Val	Gln	
	431	1		204	3	5					10					15		
	432		Thr	Glu	Asn	Lvs	Ile	Leu	Val	Lys	Gln	Ser	Pro	Leu	Leu	Tyr	Val	
	433				20	-4-				25					30	_		
	434	Asp	Ser	Asn		Val	Ser	Leu	Ser	Cys	Arg	Tyr	Ser	Tyr	Asn	Leu	Leu	
	435			35					40	_	_	_		45				
	436	Ala	Lvs	Glu	Phe	Arg	Ala	Ser	Leu	Tyr	Lys	Gly	Val	Asn	Ser	Asp	Val	
	437		50					55					60					
	438	Glu	Val	Cys	Val	Gly	Asn	Gly	Asn	Phe	Thr	Tyr	Gln	Pro	Gln	Phe	Arg	
	439	65					70					75					80	
	440	Ser	Asn	Ala	Glu	Phe	Asn	Cys	Asp	Gly	Asp	Phe	Asp	Asn	Glu	Thr	Val	
	441					85					90					95		
	442	Thr	Phe	Arg	Leu	Trp	Asn	Leu	His	Val	Asn	His	Thr	Asp	Ile	Tyr	Phe	
	443				100					105					110	_		
	444	Cys	Lys	Ile	Glu	Phe	Met	Tyr	Pro	Pro	Pro	Tyr	Leu		Asn	Glu	Arg	
	445			115					120		_			125				
	446	Ser	Asn	Gly	Thr	Ile	Ile			Lys	Glu	Lys		Leu	Cys	His	Thr	
	447		130				_	135			_	_	140	1		a 1	*** 1	
	448			Ser	Pro	Lys			Trp	Ala	Leu		vaı	vaı	ALA	GIY	val «A	
	449	145					150					155				7	160	
	450														/			
	451																	
	452						^		, *.		erd		1					
	453					,	از	1	1.1	0	Jan St.	20						
	454				NI	Son	~~	y,		\ \ \ \ \ \	<i></i>							
	455							(7									
	456																	
	457																	
	458																	

SEQUENCE VERIFICATION REPORT PATENT APPLICATION US/09/454,651

DATE: 12/20/1999 TIME: 16:47:49

INPUT SET: S34294.raw

Line	Error	Original Text
14 31 337 421	Number of Sequences (22) Doesn't Equal Actual Count (19) Wrong application Serial Number Entered (186) and Calc. Seq. Length (223) differ Entered (218) and Calc. Seq. Length (160) differ	(iii) NUMBER OF SEQUENCES: 22 (A) APPLICATION NUMBER: 08/228,208 (A) LENGTH: 186 amino acids (A) LENGTH: 218 amino acids